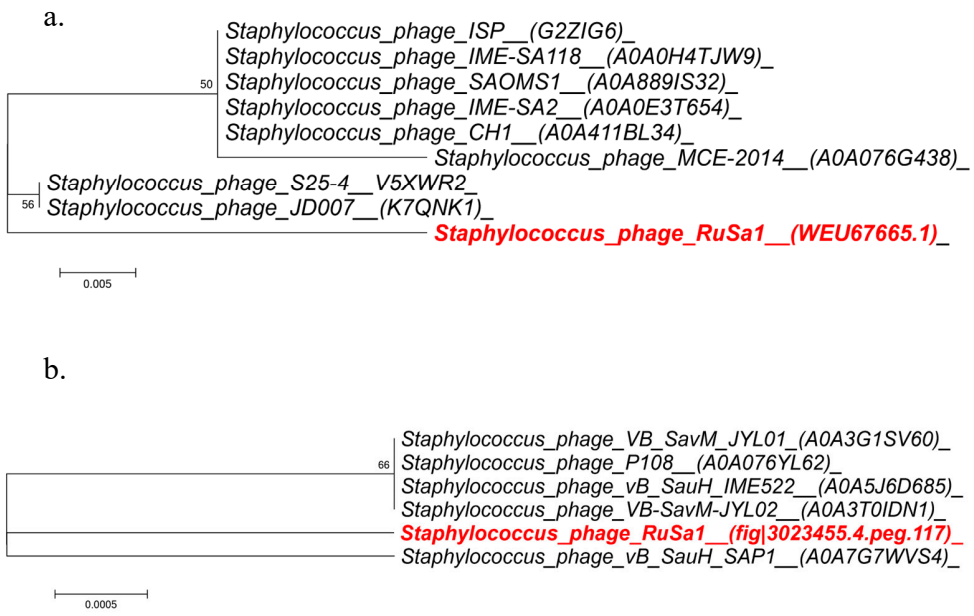


Figure S2: Phylogeny of tail proteins (a) and tail fiber proteins (b) of different *Staphylococcus* phages and RuSa1. The analysis includes different tail fiber protein and tail protein sequences retrieved from UniProt versus Phage RuSa1. The bootstrap values are indicated as branching points, with the scale bar having 0.005 (a) and 0.0005 (b) substitutions per site.